

Gene filter for polyps

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession					LC1	LC2	LC3	LC4			
Genes present in at least 4 'normal' tissues, two each upper and lower intestine, absent in polyps												
EST: mz95f11.1 Soares mouse lymph node NBMLN Mus musculus cDNA clone 721497 5', mRNA sequence. (from Genbank)	aa266897_at					538 P		A	838 P			
EST: mz46g09.r1 Barslead mouse pooled organs MRLB4 Mus musculus cDNA clone 716512 5', mRNA sequence. (from Genbank)	aa265119_s_at					523 P		A	850 P			
M.musculus gene for insulin-like growth factor binding protein-1	X67493_g_at					914 P		A	508 P			
Caspase 7	u67321_s_at					1021 P		A	457 P			
Purkinje cell protein 4	X17320_s_at					767 P		A	439 P			
EST: ms95a05.r1 Soares mouse 3NbMS Mus musculus cDNA clone 619280 5', mRNA sequence. (from Genbank)	aa175794_s_at					686 P		A	669 P			
No info for gene	ET63085_f_at					4575 P		8114 P	2117 P			
Genes present in at least 4 'polyp' tissues, two each upper and lower intestine, absent in 'normal'												

Figure 1A

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession					LC1	LC2	LC3	LC4			
EST: mz97g06.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 721402 5', mRNA sequence. [from Genbank]	AA267281.1 at											
Sim. to TRANSLATION INITIATION FACTOR EIF-2B-EPSILON SUBUNIT	Msa.8133.0 s at											
Fibroblast growth factor Inducible 16	U42385 s at											
Sim. to GLYCOGEN PHOSPHORYLASE, BRAIN FORM (EC 2.4.1.1)	Msa.6220.0 s at											
Mus musculus TRAF family member associated NF-kappa B activator (TANK) mRNA	u51907 s at											
EST: Mus musculus 3.5-dpc blastocyst cDNA 3'-end sequence, mRNA sequence. [from Genbank]	C79015.rc.a f											
Sim. to PROHIBITIN (B-CELL RECEPTOR ASSOCIATED PROTEIN 32) (BAP 32)	Msa.24245.0 f at											
EST: vb41d02.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 751491 5', mRNA sequence. [from Genbank]	AA396006 at											
Mus musculus Cdk4 and Cdk6 inhibitor p19 protein mRNA	u20497 s at											
EST: Mus musculus 3.5-dpc blastocyst cDNA 3'-end sequence, mRNA sequence. [from Genbank]	c77069.rc_s at											

Figure 1B

202210" 64T9T660

1	2	3	4	5	6	7	8	9	10	11	12	13		
Description	Accession	1	2	3	4	5	6	7	8	9	10	11	12	13
Mus musculus CDK-activating kinase assembly factor p36/MAT1	U35249_s.at	7091131P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
EST: mo35b11.r1 Life Tech mouse embryo 13.5dpc														
10666014 Mus musculus cDNA clone 555549 5', mRNA sequence. (from Genbank)	aa111610_s.at	7091131P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
EST: Mus musculus 3.5-dpc blastocyst cDNA 3'-end sequence, mRNA sequence. (from Genbank)	C81595_rc.at	7091131P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
Myosin Ia	aa285769_s.at	12021P	2162P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
EST: Mus musculus 3.5-dpc blastocyst cDNA 3'-end sequence, similar to R.norvegicus DNA sequence for LFB1/HNF1 promoter, mRNA sequence. (from Genbank)	C79518_rc.at	7091131P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
Mouse E46 mRNA for E46 protein::BRAIN PROTEIN E46	X61506_f.at	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P
EST: M.musculus expressed sequence tag M1EST/07, mRNA sequence. (from Genbank)	Z31269_s.at	13592P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P	110P

Figure 1C

202210" 64791660

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession	LC1	LC2	LC3	LC4	LC5	LC6	LC7	LC8	LC9	LC10	LC11
Sim. 16 ENDOSOMAL P24A PROTEIN PRECURSOR (70 KD ENDOMEMBRANE PROTEIN) (PHEROMONE ALPHA- FACTOR TRANSPORTER) (ACIDIC 24 KD LATE ENDOCYTIC INTERMEDIATE COMPONENT)	Msa.24715.0 _s_at											
Mus musculus mCPE-R mRNA for CPE-receptor	AB000713_g_1 _at											
Mus musculus cdc2/CDC28- like protein kinase 3 (Cik3) mRNA	AF033565_at											
Small inducible cytokine A12	u50712_s_at											
Mus musculus protein- tyrosine phosphatase mRNA	AF013490_s_1 _at											
EST: mr11h07.r1 Soares mouse 3NbMS Mus musculus cDNA clone 597181 5', mRNA sequence. (from Genbank)	aa145148_s_1 _at											
EST: mu22e08.r1 Soares 2NbMT Mus musculus cDNA clone 640166 5', mRNA sequence. (from Genbank)	aa197627_s_1 _at											
Laminin, beta 3	U43298_s_at											
Most cell protease 1	X68803_s_at											

Figure 1D

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession	1	2	3	4	5	6	7	8	9	10	11
Mus musculus non-receptor protein tyrosine kinase Ack mRNA EST: vc06102.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 765723 5', mRNA sequence, (from Genbank)	aa119775_s_at	A	A	A	A	A	A	A	A	A	A	A
Mus musculus major histocompatibility locus class III region: butyrophilin-like protein gene, partial cds; Notch4, PBX2, RAGE, lysophatidic acid acyl transferase-alpha, palmitoyl-protein thioesterase 2 (PPT2), CREB-RP, and tenascin X (TNX) genes; and CYP210HB pseudogene, complete sequence	aa277082_g_at	A	A	A	A	A	A	A	A	A	A	A
Non MHC restricted killing associated	aa270355_s_at	A	A	A	A	A	A	A	A	A	A	A
EST: vc06101.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 765762 5', mRNA sequence, (from Genbank)	L19057_s_at	A	A	A	A	A	A	A	A	A	A	A
Mus musculus C3H cytochrome P450 (Cyp1b1) mRNA	aa275330_s_at	A	A	A	A	A	A	A	A	A	A	A
Mus musculus 80kDa m-calpain subunit (calp80) mRNA	u03283_s_at	A	A	A	A	A	A	A	A	A	A	A
	d38117_s_at	A	A	A	A	A	A	A	A	A	A	A

Figure 1E

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession	LC1	LC2	LC3	LC4	LC5	LC6	LC7	LC8	LC9	LC10	LC11
EST: EST03281 Mouse 7.5 dpc embryo ectoplacental cone cDNA library Mus musculus cDNA clone C0033C02 3' similar to M.musculus VE-cadherin gene (lambda 5 clone), score = 1195, mRNA sequence. (from Genbank)	aa408463_c s_at											
EST: mc88c08.r1 Soares mouse embryo NbME13.5 14.5 Mus musculus cDNA clone 355598 5', mRNA sequence. (from Genbank)	w48388_g_at											
EST: mz98g09.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 721504 5', mRNA sequence. (from Genbank)	aa266888_s_at											
Thymidine kinase 1	m19438_s_at											
Nucleolin	X07699_s_at											
RAN GTPase activating protein 1	u08110_s_at											
Sim. to GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUB	Msa.16667.0 f_at											

Figure 1.F

1	2	3	4	5	6	7	8	9	10	11	12	13
Description	Accession					LC1	LC2	LC3	LC4			
EST: mv22d10.r1 GuayWoodford Belter mouse kidney day 0 Mus musculus cDNA clone 655795 5', mRNA sequence. (from Genbank)	aa239576_s_											
EST: m181b04.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone 636271 5', mRNA sequence. (from Genbank)	aa189313_s_											
EST: va81h09.s1 Knowles Solfer mouse 2 cell Mus musculus cDNA clone 1108769 5', mRNA sequence. (from Genbank)	aa647562_g_											
Mus musculus alpha 1 type I collagen gene, partial cds and 3' flanking region::Procollagen, type I, alpha 1	u08020_f.at											
Mouse mRNA for eptregulin IMMEDIATE EARLY PROTEIN GLY96	D30782_s.at											
EST: m131f05.r1 Soares mouse 3NbMS Mus musculus cDNA clone 599073 5', mRNA sequence. (from Genbank)	X67644_s.at											
	aa172851_s_											

Figure 1 G

14	15	16	17	18	19	20	21	22	23
UC1	UC2	UC3	UC4	Average Lower Colon	sidev Lower Colon	Average Small Intestine	sidev Small Intestine	Average Polyps	
1060 P	A	447 P	A	4	688	212.1320344	753.5	433.4564569	
539 P	A	447 P	964 P	5	686.5	231.2239174	650	275.7952139	
A	A	443 P	699 P	5	862.3333333	331.5333065	571	181.019336	
1536 P	1741 P	1883 P	957 P	6	739	398.8082246	1529.25	407.2234235	
503 P	550 P	262 P	296 P	7	593.3333333	164.8524593	402.75	144.8433061	
756 P	902 P	370 P	446 P	7	745.6666667	118.373702	618.5	252.1818127	
1780 P	1400 P	A	3096 P	7	4249.25	2818.113361	2092	890.0067415	

Figure 1 H

14	15	16	17	18	19	20	21	22	23
UC1	UC2	UC3	UC4	Average Lower Colon	stdev Lower Colon	Average Small Intestine	stdev Small Intestine	Average Polyps	
P = 104P	A	A	A	180.5	60.1040764	120.5	23.33452378	150.5	
P = 285P	A	A	A	321.5	159.0990258	303.5	20.50609665	312.5	
P = 330P	A	A	A	373	90.50966799	272	82.02438662	322.5	
P = 411P	A	A	A	456	272.9432175	271	67.88225099	363.5	
P = 390P	A	A	A	433.5	53.03300859	309	114.5512986	371.25	
P = 411P	A	A	A	414.5	79.90306627	366	253.1442277	390.25	
P = 411P	A	A	A	777.5	120.9152596	530.5	183.1406563	654	
P = 542P	A	A	A	780	677.4082964	582.5	57.27564928	681.25	
P = 542P	A	A	A	810.5	283.5498193	619.5	9.192388155	715	
P = 759P	A	A	A	1149	852.7707781	548.5	297.6919549	848.75	

Figure 1 I

14	15	16	17	18	19	20	21	22	23
	UC1	UC2	UC3	UC4	Average Lower Colon	stdev Lower Colon	Average Small Intestine	stdev Small Intestine	Average Polyps
	A	A	A	A	1259	777.8174593	554	31.11269837	906.5
	A	A	A	A	857	73.53910524	1145	236.1736649	1001
	A	A	A	A	1682	678.8225099	1932.5	1174.504364	1807.25
	A	A	A	A	5218.5	620.1326471	2819	678.8225099	4018.75
	A	A	A	A	167.3333333	51.82984983	89.5	43.13351365	136.2
	A	A	A	A	211.5	96.87362902	175.6666667	89.57864329	190
	A	A	A	A	320.5	54.44722215	397.3333333	105.8363517	366.6

Figure 1 J

14	15	16	17	18	19	20	21	22	23
	UC1	UC2	UC3	UC4	Average Lower Colon	stdev Lower Colon	Average Small Intestine	stdev Small Intestine	Average Polyps
	A	A	A	A	491.5	101.1162697	417.6666667	252.161324	447.2
	A	A	A	A	632.5	160.5132393	617	217.2487054	623.2
	A	A	A	A	727.6666667	343.7009359	833	98.99494937	769.8
	A	A	A	A	802.5	218.4959954	822	413.118627	814.2
	A	A	A	A	1035.5	161.9274529	1032.333333	367.236073	1033.6
	A	A	A	A	1349.5	4.949747468	1077.666667	436.4748943	1186.4
	A	A	A	A	1314.333333	736.7389859	1023.5	140.7142495	1198
	A	A	A	A	1495	299.5212847	919.5	979.3428919	1264.8
	A	A	A	A	2416	516.1879503	1071	683.2854455	1609

Figure 1, K

2002101547650

	14	15	16	17	18	19	20	21	22	23
	UC1	UC2	UC3	UC4		Average Lower Colon	stddev Lower Colon	Average Small Intestine	stddev Small Intestine	Average Polyps
P12412P		A	A	A	A 5	1966.5	760.1397898	1761.333333	883.1224906	1843.4
P12413P										
P12414P	A	A	A	A	A 5	3350	69.29646456	1780.333333	193.8874244	2408.2
P12415P										
P12416P										
P12417P	A	A	A	A	A 5	5575.666667	2866.201726	4153.5	614.4757929	5006.8
P12418P	A	A	A	A	A 6	133	9.539392014	221.6666667	100.4506512	177.3333333
P12419P										
P12420P	A	A	A	A	A 6	551	175.8379936	458.3333333	254.4097744	504.6666667
P12421P										
P12422P	A	A	A	A	A 6	725.5	78.48885271	414.5	150.529067	518.1666667
P12423P										
P12424P	A	A	A	A	A 6	715	845.6997103	433.5	47.0496192	527.3333333

Figure 4L

14	15	16	17	18	19	20	21	22	23
UC1	UC2	UC3	UC4	Average Lower Colon	stdev Lower Colon	Average Small Intestine	stdev Small Intestine	Average Polyps	
P-0101B	A	A	A	A	760.5	92.6309834	741	220.5190846	747.5
P-0102B	A	A	A	A	1070	644.8813844	753	368.3150825	858.6666667
P-0103B	A	A	A	A	1483	472.8435259	680.3333333	105.9874206	1081.666667
P-0104B	A	A	A	A	1658.666667	532.3244625	1183.666667	209.8340614	1421.166667
P-0105B	A	A	A	A	2394.666667	523.1102497	1965.333333	683.7648231	2180
P-0106B	A	A	A	A	3473	813.9428727	1345	1014.261801	2409
P-0107B	A	A	A	A	3706	1114.400287	2582.5	288.1301789	2957

Figure 1 M

202210" 64T9T660

14	15	16	17	18	19	20	21	22	23
	UC1	UC2	UC3	UC4	Average Lower Colon	stdev Lower Colon	Average Small Intestine	stdev Small Intestine	Average Polyps
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 7	250.6666667	276.5	77.71100308	265.4285714
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 7	87.70784838	276	77.78817391	314.2857143
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 7	266.642551	606.6666667	277.5776168	953.5714286
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 7	328.82267021	927	745.4828413	1056.428571
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 8	95.47600047	307.25	72.49080401	338.875
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 8	438.9605145	351.75	149.0936507	469.125
202210" 64T9T660									
202210" 64T9T660	A	A	A	A	A 8	750.392786	2223.25	438.7720555	2621.625

Figure 1N

24	25	26	27
stddev polyps	Average Normal Tissue	stddev Normal Tissue	Description
	720.75	281.1729895	Similar to Rat Oplold growth factor receptor, which regulates cellular renewal, wound healing. OGF inhibits pancreatic and squamous cell carcinomas
	664.6	227.5902019	
	745.8	297.6771741	Insulin-like growth factor binding protein 1, high affinity, expressed in liver, decidua, kidney and in amniotic fluid, regulator of apoptosis
	1265.833333	545.7473469	caspase 7, effector, cysteine containing aspartate-specific protease, CASP3 subfamily, stored in the mitochondrial intermembrane space and released into cytosol after appropriate apoptotic stimuli, promoting apoptosis, H3 isoform alpha, interacting with calpain during B cell clonal deletion by apoptosis
	484.4285714	172.9921551	Purkinje cell protein 4, rat PEP-19, neuron-specific polypeptide homolog gene, with homology to S100 calcium binding proteins, involved in the development of the central nervous system
	673	202.704218	
	3324.714286	2358.930952	

Figure 10

24 stdev polyps	25 Average Normal Tissue	26 stdev Normal Tissue	27 Description
50.849451			
93.196924			eukariotic translation initiation factor 2B, subunit 5, ubiquitously expressed (epsilon, 82kDa)
91.507741			
194.36306			phosphorylase, glycogen catabolism, brain
102.36332			tumor necrosis receptor-associated factor, TRAF-interacting protein, mediator of NFkB activation after induction by TRAF2, apoptosis inhibitor?
155.79768			
190.76163			prohibitin (antiproliferative protein) potential tumor suppressor regulating E2F1 function
408.72515			
197.45548			cyclin dependent kinase 4 and 6 (CDK4/CDK6) inhibitor, p19, regulator of the cell cycle, passage through the G1 checkpoint, expressed primarily in hematopoietic tissues
626.21582			

Figure 1P

2022101647910179

24	25	26	27
sidev polyps	Average Normal Tissue	sidev Normal Tissue	Description
606.35331			
219.1879			
796.45271			
1483.5728			myosin IC, unconventional, apparently non filamentous, homologous to mouse Myo1e, crypt cell marker
60.213786			
82.118816			
90.071083			

Figure 1 Q

24	25	26	27
std dev polyps	Average Normal Tissue	std dev Normal Tissue	Description
189.69502			carboxypeptidase E, metallo carboxypeptidase family, regulated secretory pathway sorting receptor, involved in the timing of paired basic residues at the C terminus of prohormone-derived peptides, mutated in the obese fat mouse
173.52723			CDC-like kinase 3, with two alternatively spliced forms, one catalytically active and one inactive isoforms, interacting with and inducing the nuclear redistribution of SR proteins SFRS* (see symbols), widely expressed
254.64426			
312.06201			
272.00974			
342.67959			
549.28818			laminin 5 (kalinin/ricin), beta 3 polypeptide, component of cutaneous basement membrane zone, expressed in stratifying squamous epithelium, downstream from E-cadherin
619.67306			chymase, mast cell, with a variant putatively involved in eczema
918.01934			

Figure 1R

24	25	26	27
sidev polyps	Average Normal Tissue	sidev Normal Tissue	Description
739.61733			Inhibits Ras-induced malignant phenotypes in fibroblasts ?? http://www.kfinder.com/member-search/getdoc.cgi?ord=5&searchId=1&have_local_holdings_file=1&local_journals_only=0
871.29369			
2192.8791			
80.193932			
202.07292			cytochrome p450, family 1 (aromatic compound inducible), member B1, expressed in ocular structures of the anterior uveal tract, possibly involved in the metabolism of substances active in the eye growth and differentiation
201.54346			calpain, large polypeptide 11, calcium dependent neutral cysteine proteinase, papain superfamily, mu type (not mutated in MEN1)
406.81822			

Figure 1S

24	25	26	27
stdev polyps	Average Normal Tissue	stdev Normal Tissue	Description
176.05312			In different types of epithelial tumours, cadherin expression is inversely correlated with invasiveness and metastatic dissemination
437.45293			hGAR1 is a component of HJACA snRNPs and telomerase in vivo
535.91778			Mus musculus cleavage and polyadenylation specificity factor 73 kDa subunit mRNA
445.69874			thymidine kinase 1, pyrimidine salvage pathway, soluble, putative up-regulated c-Myc target gene, nucleolin, major multifunctional nucleolar protein of exponentially growing cells, characterized by unique tripartite function within each domain, performing activities i.e. a specific DNA helicase and DNA-dependent ATPase, also acting as a sequence-specific RNA binding protein an autoantigen, a component of B cell specific transcription factor, involved in ribosome biogenesis, cytokinesis, nucleogenesis, cell proliferation and growth, chromatin remodeling etc..
593.10168			Inhibits gtp exchange on ran. forms a ran-gtp-ranbp1 trimeric complex. Increase gtp hydrolysis induced by the ran gtpase activating protein rangap1. may act in an intracellular signaling pathway which may control the progression through the cell cycle by regulating the transport of protein and nucleic acids across the nuclear membrane.
1425.5368			guanine nucleotide-binding proteins (g proteins) are involved as a modulator or transducer in various transmembrane signaling systems. the beta and gamma chains are required for the gtpase activity, for replacement of gdp by gtp, and for g protein- effector interaction.
796.73684			

Figure 1 T

24	25	26	27
stdev polyps	Average Normal Tissue	stdev Normal Tissue	Description
68.397995			ubiquitin conjugating enzyme E2 variant 1, expressed as at least four isoforms, transcriptional activator of FOS promoter, underexpressed in hormone refractory prostate cancer, potentially involved in the control of differentiation and the entry of a larger proportion of cells in the division cycle and an accumulation in G2-M
84.533454			
408.08367			
583.072			collagen type I, alpha 1, fibril forming, putative downregulated c-Myc target gene, http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&list_uids=10769639&dopt=Abstract Increased expression of heparin binding EGF (HB-EGF), amphiregulin, TGF alpha and epiregulin in androgen-independent prostate cancer cell lines. PRG1: a novel early-response gene transcriptionally induced by pituitary adenylate cyclase activating polypeptide in a pancreatic carcinoma cell line.
85.4508			
328.4076			
710.77944			

Figure 1U

FIGURE 1V

1A	1H	1O
1B	1I	1P
1C	1J	1Q
1D	1K	1R
1E	1L	1S
1F	1M	1T
1G	1N	1U